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RAW SEQUENCE LISTING DATE: 09/07/2001
PATENT APPLICATION: US/09/840,277 TIME: 14:29:07

Input Set : A:\A-688A.ST25.txt
Output Set: N:\CRF3\09072001\I840277.raw

Does not have a statistics sheet

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p. 5

```
  3 <110> APPLICANT: FEIGE, ULRICH
  4       KOHNO, TADAHIKO
  5       LACEY, DAVID LEE
  6       BOONE, THOMAS CHARLES
  8 <120> TITLE OF INVENTION: INTEGRIN/ADHESION ANTAGONISTS
 10 <130> FILE REFERENCE: A-688A
 12 <140> CURRENT APPLICATION NUMBER: 09/840,277
C--> 13 <141> CURRENT FILING DATE: 2001-08-14
 15 <150> PRIOR APPLICATION NUMBER: 60/198,919
 16 <151> PRIOR FILING DATE: 2000-04-21
 18 <150> PRIOR APPLICATION NUMBER: 60/201,394
 19 <151> PRIOR FILING DATE: 2000-05-03
 21 <160> NUMBER OF SEQ ID NOS: 135
 23 <170> SOFTWARE: PatentIn version 3.1
 25 <210> SEQ ID NO: 1
 26 <211> LENGTH: 684
 27 <212> TYPE: DNA
 28 <213> ORGANISM: Homo sapiens
 30 <220> FEATURE:
 31 <221> NAME/KEY: CDS
 32 <222> LOCATION: (1)..(684)
 33 <223> OTHER INFORMATION:
 36 <400> SEQUENCE: 1
 37 atg gac aaa act cac aca tgt cca cct tgt cca gct ccg gaa ctc ctg       48
 38 Met Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu
 39 1               5                  10                  15
 41 ggg gga ccg tca gtc ttc ctc ttc ccc cca aaa ccc aag gac acc ctc       96
 42 Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu
 43             20                  25                  30
 45 atg atc tcc cgg acc cct gag gtc aca tgc gtg gtg gtg gac gtg agc      144
 46 Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser
 47         35                  40                  45
 49 cac gaa gac cct gag gtc aag ttc aac tgg tac gtg gac ggc gtg gag      192
 50 His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu
 51     50                  55                  60
 53 gtg cat aat gcc aag aca aag ccg cgg gag gag cag tac aac agc acg      240
 54 Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr
 55 65                  70                  75                  80
 57 tac cgt gtg gtc agc gtc ctc acc gtc ctg cac cag gac tgg ctg aat      288
 58 Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn
 59                 85                  90                  95
 61 ggc aag gag tac aag tgc aag gtc tcc aac aaa gcc ctc cca gcc ccc      336
 62 Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro
 63             100                 105                 110
 65 atc gag aaa acc atc tcc aaa gcc aaa ggg cag ccc cga gaa cca cag      384
 66 Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln
 67         115                 120                 125
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69 gtg tac acc ctg ccc cca tcc cgg gat gag ctg acc aag aac cag gtc      432
70 Val Tyr Thr Leu Pro Pro Ser Arg Asp Glu Leu Thr Lys Asn Gln Val
71     130                 135                 140
73 agc ctg acc tgc ctg gtc aaa ggc ttc tat ccc agc gac atc gcc gtg      480
74 Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val
75 145                 150                 155                 160
77 gag tgg gag agc aat ggg cag ccg gag aac aac tac aag acc acg cct      528
78 Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro
79                 165                 170                 175
81 ccc gtg ctg gac tcc gac ggc tcc ttc ttc ctc tac agc aag ctc acc      576
82 Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr
83             180                 185                 190
85 gtg gac aag agc agg tgg cag cag ggg aac gtc ttc tca tgc tcc gtg      624
86 Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val
87         195                 200                 205
89 atg cat gag gct ctg cac aac cac tac acg cag aag agc ctc tcc ctg      672
90 Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu
91     210                 215                 220
93 tct ccg ggt aaa                                                    684
94 Ser Pro Gly Lys
95 225
98 <210> SEQ ID NO: 2
99 <211> LENGTH: 228
100 <212> TYPE: PRT
101 <213> ORGANISM: Homo sapiens
103 <400> SEQUENCE: 2
105 Met Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu
106 1               5                  10                  15
109 Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu
110             20                  25                  30
113 Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser
114         35                  40                  45
117 His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu
118     50                  55                  60
121 Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr
122 65                  70                  75                  80
125 Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn
126                 85                  90                  95
129 Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro
130             100                 105                 110
133 Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln
134         115                 120                 125
137 Val Tyr Thr Leu Pro Pro Ser Arg Asp Glu Leu Thr Lys Asn Gln Val
138     130                 135                 140
141 Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val
142 145                 150                 155                 160
145 Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro
146                 165                 170                 175
149 Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr
```

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PATENT APPLICATION: US/09/840,277 TIME: 14:29:07

Input Set : A:\A-688A.ST25.txt
Output Set: N:\CRF3\09072001\I840277.raw

```
150             180                 185                 190
153 Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val
154         195                 200                 205
157 Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu
158     210                 215                 220
161 Ser Pro Gly Lys
162 225
165 <210> SEQ ID NO: 3
166 <211> LENGTH: 8
167 <212> TYPE: PRT
168 <213> ORGANISM: Artificial Sequence
170 <220> FEATURE:
171 <223> OTHER INFORMATION: Preferred linker
173 <400> SEQUENCE: 3
175 Gly Gly Gly Lys Gly Gly Gly Gly
176 1               5
179 <210> SEQ ID NO: 4
180 <211> LENGTH: 8
181 <212> TYPE: PRT
182 <213> ORGANISM: Artificial Sequence
184 <220> FEATURE:
185 <223> OTHER INFORMATION: Preferred linker
187 <400> SEQUENCE: 4
189 Gly Gly Gly Asn Gly Ser Gly Gly
190 1               5
193 <210> SEQ ID NO: 5
194 <211> LENGTH: 8
195 <212> TYPE: PRT
196 <213> ORGANISM: Artificial Sequence
198 <220> FEATURE:
199 <223> OTHER INFORMATION: Preferred linker
201 <400> SEQUENCE: 5
203 Gly Gly Gly Cys Gly Gly Gly Gly
204 1               5
207 <210> SEQ ID NO: 6
208 <211> LENGTH: 5
209 <212> TYPE: PRT
210 <213> ORGANISM: Artificial Sequence
212 <220> FEATURE:
213 <223> OTHER INFORMATION: Preferred linker
215 <400> SEQUENCE: 6
217 Gly Pro Asn Gly Gly
218 1               5
221 <210> SEQ ID NO: 7
222 <211> LENGTH: 5
223 <212> TYPE: PRT
224 <213> ORGANISM: Artificial Sequence
226 <220> FEATURE:
227 <223> OTHER INFORMATION: Laminin peptide
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RAW SEQUENCE LISTING DATE: 09/07/2001
PATENT APPLICATION: US/09/840,277 TIME: 14:29:07

Input Set : A:\A-688A.ST25.txt
Output Set: N:\CRF3\09072001\I840277.raw

```
229 <400> SEQUENCE: 7
231 Tyr Ile Gly Ser Arg
232 1               5
235 <210> SEQ ID NO: 8
236 <211> LENGTH: 49
237 <212> TYPE: PRT
238 <213> ORGANISM: Artificial Sequence
240 <220> FEATURE:
241 <223> OTHER INFORMATION: Echistatin peptide
243 <400> SEQUENCE: 8
245 Glu Cys Glu Ser Gly Pro Cys Cys Arg Asn Cys Lys Phe Leu Lys Glu
246 1               5                   10                  15
249 Gly Thr Ile Cys Lys Arg Ala Arg Gly Asp Asp Met Asp Asp Tyr Cys
250             20                  25                  30
253 Asn Gly Lys Thr Cys Asp Cys Pro Arg Asn Pro His Lys Gly Pro Ala
254         35                  40                  45
257 Thr
261 <210> SEQ ID NO: 9
262 <211> LENGTH: 7
263 <212> TYPE: PRT
264 <213> ORGANISM: Artificial Sequence
266 <220> FEATURE:
267 <223> OTHER INFORMATION: RGD, NGR derivative peptide
269 <220> FEATURE:
270 <221> NAME/KEY: misc_feature
271 <222> LOCATION: (2, 5 and)..(7)
272 <223> OTHER INFORMATION: Xaa is any amino acid
275 <400> SEQUENCE: 9
W--> 277 Arg Xaa Glu Thr Xaa Trp Xaa
278 1               5
281 <210> SEQ ID NO: 10
282 <211> LENGTH: 7
283 <212> TYPE: PRT
284 <213> ORGANISM: Artificial Sequence
286 <220> FEATURE:
287 <223> OTHER INFORMATION: RGD, NGR derivative peptide
289 <220> FEATURE:
290 <221> NAME/KEY: misc_feature
291 <222> LOCATION: (2, 5 and)..(7)
292 <223> OTHER INFORMATION: Xaa is any amino acid
295 <400> SEQUENCE: 10
W--> 297 Arg Xaa Glu Thr Xaa Trp Xaa
298 1               5
301 <210> SEQ ID NO: 11
302 <211> LENGTH: 9
303 <212> TYPE: PRT
304 <213> ORGANISM: Artificial Sequence
306 <220> FEATURE:
307 <223> OTHER INFORMATION: RGD, NGR derivative peptide
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RAW SEQUENCE LISTING DATE: 09/07/2001
PATENT APPLICATION: US/09/840,277 TIME: 14:29:07

Input Set : A:\A-688A.ST25.txt
Output Set: N:\CRF3\09072001\I840277.raw

```
     309 <220> FEATURE:
     310 <221> NAME/KEY: misc_feature
     311 <222> LOCATION: (2, 3, 7 and)..(8)
     312 <223> OTHER INFORMATION: Xaa is any amino acid
     315 <400> SEQUENCE: 11
W--> 317 Cys Xaa Xaa Arg Leu Asp Xaa Xaa Cys
     318 1               5
     321 <210> SEQ ID NO: 12
     322 <211> LENGTH: 7
     323 <212> TYPE: PRT
     324 <213> ORGANISM: Artificial Sequence
     326 <220> FEATURE:
     327 <223> OTHER INFORMATION: RGD, NGR derivative peptide
     329 <220> FEATURE:
     330 <221> NAME/KEY: misc_feature
     331 <222> LOCATION: (2 and)..(3)
     332 <223> OTHER INFORMATION: Xaa is any amino acid
     335 <400> SEQUENCE: 12
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     338 1               5
     341 <210> SEQ ID NO: 13
     342 <211> LENGTH: 9
     343 <212> TYPE: PRT
     344 <213> ORGANISM: Artificial Sequence
     346 <220> FEATURE:
     347 <223> OTHER INFORMATION: RGD, NGR derivative peptide
     349 <220> FEATURE:
     350 <221> NAME/KEY: misc_feature
     351 <222> LOCATION: (1, 2, 3, 7, 8 and)..(9)
     352 <223> OTHER INFORMATION: Xaa is any amino acid
     355 <400> SEQUENCE: 13
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     358 1               5
     361 <210> SEQ ID NO: 14
     362 <211> LENGTH: 9
     363 <212> TYPE: PRT
     364 <213> ORGANISM: Artificial Sequence
     366 <220> FEATURE:
     367 <223> OTHER INFORMATION: RGD, NGR derivative peptide
     369 <220> FEATURE:
     370 <221> NAME/KEY: misc_feature
     371 <222> LOCATION: (2 )..(8)
     372 <223> OTHER INFORMATION: Xaa is any amino acid
     376 <400> SEQUENCE: 14
W--> 378 Cys Xaa Cys Arg Gly Asp Cys Xaa Cys
     379 1               5
     382 <210> SEQ ID NO: 15
     383 <211> LENGTH: 8
     384 <212> TYPE: PRT
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Use of n and / or Xaa has been detected in the Sequence Listing. Review the Sequence Listing to ensure a corresponding explanation is present in the <220> to <223> fields of each sequence using n or Xaa.

VERIFICATION SUMMARY DATE: 09/07/2001
PATENT APPLICATION: US/09/840,277 TIME: 14:29:08

Input Set : A:\A-688A.ST25.txt
Output Set: N:\CRF3\09072001\I840277.raw

L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:277 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:297 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:317 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:337 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:357 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:378 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
L:410 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:448 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16
L:468 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:492 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18
L:512 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19
L:516 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19
L:536 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20
L:556 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21
L:828 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:40
L:974 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:50
L:1120 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:58
L:1140 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:59
L:1536 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:86
L:1556 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:87